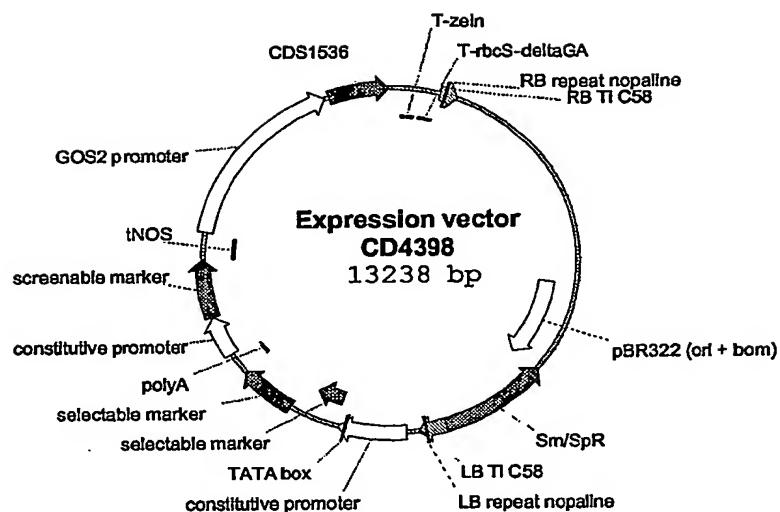
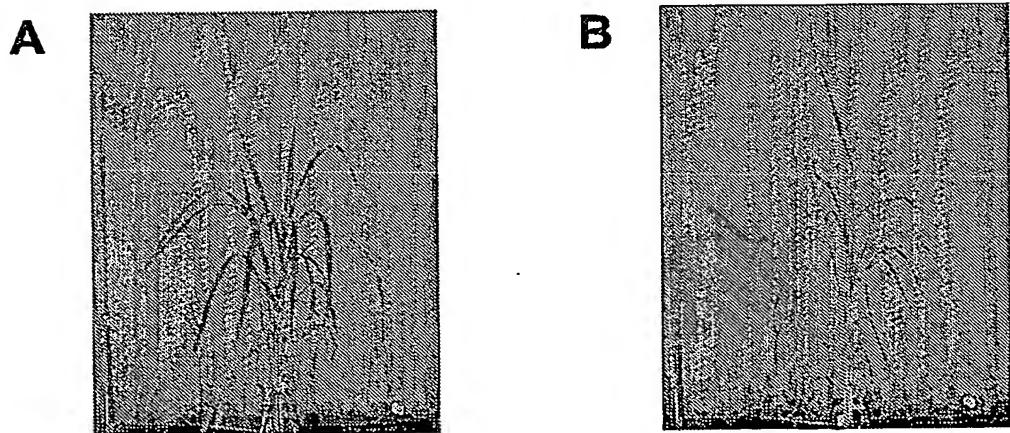


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**FIGURE 1****FIGURE 2**

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SEQ ID NO 1: *Arabidopsis thaliana* STZ cDNA (CDS1536)

AATGGCGCTCGAGGCTCTTACATCACCAAGATTAGCTCTCCGATTCCTCTTGTTCGAAG
ATTCTCAGTCTTCCATGGAGTCGAGCACTGGACAAAGGGTAAGCGATCTAAGAGATCAAGA
TCCGATTCCACCAACCAAAACCTCACTGAGGAAGAGTATCTAGCTTTGCCTCATGCTTCT
CGCTCGCGACAACCGTCAGCCTCCTCCCTCCGGCGGTGGAGAAGTTGAGCTACAAGTGA
GCGTCTGCAGACAAGACGTTCTTCAAGCTCTCGGTGGTACAAGGCAAGGCCACCGT
AAGAACTTATCACAGACTCTCCGGCGGAGGAGATGATCATTCAACCTCGTCGGCGACAAC
CACATCCGCCGTGACTACTGGAAGTGGAAATCACACGTTGCACCATCTGTAACAAGTCTT
TTCCTCCGGTCAAGCTCTCGGCGACACAAGCGGTGCACTACGAAGGAAACAACACATC
AACACTAGTAGCGTGTCCAACCTCGAAGGTGCGGGGTCCACTAGCCACGTTAGCAGTAGCCA
CCGTGGGTTTGACCTAACATCCCTCCGATCCCTGAATTCTCGATGGTCAACGGAGACGACG
AAGTCATGAGCCCTATGCCGGCGAAGAAGCCTCGGTTGACTTCCGGTCAAACTTCAACTT
****TAAGGAAATT****

SEQ ID NO 2: *Arabidopsis thaliana* STZ protein with annotation of the domains

B-BOX	L-BOX
MALEALTSPRLASPIPPLFEDSSVFHGVEHWTKG KRSKRSRS DFHHQNLTEEYLA FCLMLL	
ARDNRQPPP PPPAVEKLSYK CSVCDKTFSSYQALGGHKASHRKNL SQTLGGDDHSTSSATT	
TSAVTGS GKSHV CTICNKSFP SGQALGGHKRCH Y EGNNNNINTSSVSNSEGAGSTSHVSSSH	
RGFD LNLIPPIPEFSMVNGD EVMSPM PAKPR DFPVKLQL	

EAR motif

SEQ ID NO 3: PRM3204 (sense, start codon in italics)

5' **GGGGACAAGTTGTACAAAAAAGCAGGCTTCACAATGGCGCTCGAGGCTC** 3'

SEQ ID NO 4: PRM3205 (reverse, complementary stop codon in italics)

5' **GGGGACCAC~~TTGTACAAGAAAGCTGGGTA~~ATTCC~~TTAAAGTTGAAGTTGA~~** 3'

BOXES**SEQ ID NO 5: QALGGH motif**

QALGGH

SEQ ID NO 6: NNM motif

NN (M/W) QMH

SEQ ID NO 7: EAR motif

hDLNh (X) P

SEQ ID NO 8: B-Box

KR (S) KRXR

SEQ ID NO 9: L-Box

EXEXXAXCLXXL

FIGURE 3

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ORTHOLOGS OF STZ in OTHER PLANT SPECIES

SEQ ID NO 10: Dg_AF119050_1 *Datisca glomerata* zinc-finger protein 1 (zfp1) mRNA, complete cds

GGCACGAGGACAAATTCTCTCTATCCTCTGAATATCTTGGTTGTGAACGTGAGAACGTA
 TTAGATGGCTCTAGAACGCGCTCAACTCTCCGACCACAGCTACGCCGGTGTTCACTACCGACG
 ACCCCAGCTGAATTACCTTGAGCCATGGACCAAGCGTAAGCGTCCAAAGCGTACGCCGCTTA
 GATAGCCCCATAACCGAGGAAGAGTACCTTGCTTCTGCCTCATCATGCTCGCTCGTGGCCGC
 GTTGCCTCTGCAAATCGACGGGATTCTCAGTCTTCATTAGTACGCCGTGAAGCAACGAC
 TTCGGCTACCAAAGTCAGTTATAAGTGCTCTGTGCGATAAGGCCCTTCTGTCTATCAGG
 CTTGGGTGGGCACAAGGCCAGCCACAGAAAGCTCGCTGGCGGAAGATCAATCGACCTCC
 TTTGCCACCACGAATTCAAGCCACCGTCACTACCACACGCCCTCCGGAGGTGGTGGCAGGTC
 TCATGAGTGTCTATTGCCACAAATCGTCCCGACTGCCAGGCCTGGGTGGTCACAAGC
 GCTGCCACTACGAAGGCAGTATCGCGGCAATAGTATTACCAACCACAACACCACAAAC
 AGCGGAAGCAACGGTGGCATGAGCATGACCTCGAAGTAGGTTCCACACACAGTCAGCCA
 CAGTCACCGTGACTTCGATCTCAACATCCCGGCTTGCCGGAGTTCGGTGAATTCTCA
 TATCCGGGGATGACGAGGTGAGAGTCCTCATCCGGCCAAGAAACCCGTATATTGATGAAA
AAAAACATTCTCAAGATCACTGAACCAGGCTTAGTTCTTATAGGAGGAGATTAAAAAA
 AGTAGTATCTCTTTCTTATCCGTAGGATAATTAATATTCGTGTACATAAAATTGTA
 GTTCTTAACACACTCTGTTCACTTGTGCTCAACTTGTATTGGTTATTCATT
 ATGAAAATTCAATT

SEQ ID NO 11: Dg_AF119050_1 *Datisca glomerata*, STZ ortholog, protein

MALEALNSPTTATPVFHDDPSLNYLEPWTKRKRSKRTRLDSPHTEEYLAFCILMLARGRV
 ASANRDSQSSIQIPEATTSATKVSYKCSVCDKAFFSYQALGGHKASHRKLAGEDQSTS
 ATTNSATVTTTASGGGGRSHECSICHKSFTPQALGGHKRCHYEGSIGGNSIHHHNNTNS
 GSNGGMSMTSEVGSTHTVSHSHRDFDLNIPALPEFRSNFFISGDDEVESPHPAKKPRILMK

SEQ ID NO 12: Gm_T09602_U68763.1_GMU68763 Glycine max (soybean) probable zinc finger protein SCOF-1 mRNA, complete cds

AAAATTCTCACTCTCTCATCTCGAGATCATAGTATCATATTCAATATCATTTCATACC
 AAACACATGGCTTGGAAAGCTCTCAACTCACCAACAACACCCTCCATCTTCCCTTGA
 CGACCCAACATTCCATGGCGAACGAAAACGTTCAAAGCGTTCTCGGACCATCCTCTG
 AAGAAGAGTACCTCGCCCTCTGCCTCATCATGCTCGCTCGCGGCGACCACCCACCGTCAAC
 AACCGCCACGTCAGCCCTCCGCCGCTACAGCCACAGCCACAGCCACACCAGATCCTCCAC
 CAAGCTCAGTTACAATGCTCCGTTGCGACAAGAGCTCCCTCTTACCAAGCGCTCGGTG
 GACACAAAGGCCAGTCACCGGAAACTCGCCGGCCGCCGAAGACCAACCCCCCAGCACC
 ACTTCTCCGCCGCCACAGCTCCGCTCCGGAGGTTAAGGCCCAGTGCTCCATTG
 CCACAAATCCTCCCCACCGGACAGGCCCTGGCGGACACAAACGTTGTCACTACGAAGGTA
 ACGGTAACGAAATAACAACAAGTAACAGCGTTGTCAACCGTCGCCTCGGAAGGCCTGGC
 TCCACCCACACTGTCAGTCACGCCACCCACCGCAGCTCGATCTCAACATCCGGCTTCC
 GGATTTCGACCAAGGTGGAGAACGAGCAGGTTGAGAGGCCCTCACCTGTCATGAAGAAC
 CTCGCCTCTCGTCATTCCAAGATCGAAATCCCCAATTCAATGAACACTCGTTGAATT
 AGTTTATTTTCGACTATATTTGGAGAATTGAGAGTTACTATAATTGATTGTTAC
 ATAGTACTTGGAAAGTTGTTGGACCGTACCGGACCCAGTTCTGGTTGAGGTTGTACTTT
 CACAACAGTGGCAGATTGCAATTCAATTATTGTTATTAAAAAA
 AAAA

FIGURE 3 (continued)

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SEQ ID NO 13: Gm_T09602 Glycine max (soybean) probable zinc finger protein SCOF-1, STZ ortholog, protein
 MALEALNSPTTAPSFPFDDPTIPWAKRKRSKRSRDHPSEEEYLALCLIMLARGGTTVNNR
 HVSPPPLQPQPQPTPDPSTKLSYKCSVCDKSFPSYQALGGHKASHRKLAGAAEDQPPSTTS
 SAAATSSASGGKAHECSICHKSFTPQALGGHKRCHYEGNGNGNNNNNSNVTVASEGVGST
 HTVSHGHHRDFDLNIPAFPDFSTKVGDEVESPHPVMKKPRLFVIPKIEIPQFQ

SEQ ID NO 14: Ms_CAB77055_Y18788.1_MSY18788 Medicago sativa putative TFIIIA (or kruppel)-like zinc finger protein mRNA
 AATTCCGGCACGAGAAATAACCACTCTCTCTCAAAACCTCCTTGCCCTTGCTTCACTT
 TCACTTGCCTAACGCTAACACTCTCTCGAGTGTCTTCACTCATATGGCTATGGA
 AGCACTTAACCTACCCACCCTGCTACTCCTTCACACCCCTTGAGGAACCAAATCTGAGTT
 ATCTTGAAACACCGTGGACGAAAGGTAACGATCAAAGCGTCTCGCATGGATCAATCTCA
 TGCACCTGAAGAAGAGTATCTCGCTTTGTCTCATCATGCTTGCTCGCAGCGGTAAACAACAA
 CGACAAAAAGTCTGATTGGTGGCGACGCCGCTAACACCCTTAAACTCAGTCACAAATGCT
 CAGTCTGCAACAAAGCTTCTCATCTTATCAAGCCCTAGGTGGACACAAAGCCAGTCACCGG
 AAAGCTGTTATGTCCGCAACCACCGCTGAAGATCAGATCACCACCTCATCCGCCGTGAC
 TACCAAGCTCTGCTTCAAACGTAAGAACACAAGACTCATGAGTGTCCATCTGTCACAAATCCT
 TCCCTACTGGACAGGCTTGGGAGGACACAAGCGTTGTCACTACGAAGGCAGCGTTGGGCC
 GGTGCCGGTGCTGGAAAGTAACGCTGTAACTCGCTCTGAAGGGATTGGATTGTACACAGCA
 CCACCGTGAATTGATCTAACCTCCGGCTTCCGGACTTTCAAAGAAGTTTGTGG
 ATGACGAGGTTTTAGTCCTTACCTGCTGCAAAGAACGCCCTGTCTTCAAAGCTGGAAATT
 CCTTCTCATTACTGATCAATAATAGATCCAATTTTATTGTTATTATTAAATAATTATT
 CGCTTAGGGCATAGTTATTCTTCAATTATTGATCAATTGTTCTGTACA
 TACAAATTGGGATTGGTTAGAATTAGGACGGTTGTAGACAATGGAAATTCAATTCAATT
 ATTTAATTGTGT

SEQ ID NO 15: Ms_CAB77055_Medicago sativa putative TFIIIA (or kruppel)-like zinc finger protein, STZ ortholog, protein
 MAMEALNSPTTATPFTPFEENLSELETPWTKRKRSKRSRMDQSSCTEEYLALCLIMLARS
 GNNNDKKSDSVATPLTTVKLSHKCSVNCNAFKSSYQALGGHKASHRKAVMSATTAEDQITTS
 SAVTTSSASNGKNKTHECSICHKSFTPQALGGHKRCHYEGSVGAGAGAGSNAVTASEGVGL
 SHSHHRDFDLNLPAPDFSKKFFVDEVFSPLPAAKPCLFKLEIPSHY

SEQ ID NO 16: Nt_AAC06243_AF053077 Nicotiana tabacum osmotic stress-induced zinc-finger protein (zfp) mRNA, complete cds
 TTTTCCCTGAATTGATAACTAAAGAGAAATTATGACTCTTGAAGCTTGAAAGTCACCTA
 CGGCCGCAACGCCGACTCTACCACACCGTATGAAGATGATGATGAAATTCATAATTGGAT
 TCTTGGGCTAAAGAAAACGATCAAAACGGCCCCGTATTGATGCCCAACCGACTGAAGAAGA
 GTATTAGCCCTCTGTCTCATCATGCTCGCTCGCAGCGGAACCGGAACCAGAACCGGTTAA
 CTGATGCTACTACTTCCCAACAACCTGCCATAAAAAACGCCAGTTGCCCGGTTCAT
 AAGAAAGAGGTGGCAACAGAGCAAGCAGAGCAATCTTACAAGTGTAGCGTGTGTGACAAGGC
 TTTTCTTCTTATCAAGCACTCGGTGGGCATAAAAGCAAGTCACCGTAAACTACTACTAG
 CTACCGCCGCTCTGATGATAACAATCCTCAACTTCAACTTCCACTGGCGCCGTTAATATC
 TCTGCTCTTAATCCAACCTGGTCGTTCACACGCTCTGTTCTATTGCCACAAGCTTTCTAC
 TGGCCAAGCTTGGGTGGCACAAGCGCCGCACTATGAAGGCAAACTCGGTGGTAACAGCC
 GCGACTTAGGCCGGCGCGGGCGGGTCAAGTGGAAAGCGTCTGACTACTCAGACGGC

FIGURE 3 (continued)

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GGCGCGTCGACTCACACGCTACGTGACTTTGACCTAACATGCCTGCTCGCCGGAATTGCA
 ACTGGGTCTGAGTATTGATTGTGGACGGAAAAGTCACACTGTTGCCATGGTCCAAGAGGTGG
 AAAGTCCTATGCTGCAAAGAACCGCGTTATTGTTCTGGGTTGAAACTTCTTAGG
 GGAATTGAATTGATTGTGTTAGCCAAATTAGTAAATTGGTCATGTGATTTTATTTAG
 GAAAAGGAATTATTGATTGTTACCGTTATTCTTAGGGTGGTATTATGTACAGGGAGTG
 AATCATTCAATTGTTTACACTTCTTAATTATATATTCTTTTTACACATAAAAAAAA
 AAAAAAA

SEQ ID NO 17: Nt_AAC06243_Nicotiana tabacum osmotic stress-induced zinc-finger protein, STZ ortholog, protein
 MTLEALKSPTAATPTLPPRYEDDEIHNLDWSAKGKRSPRIDAPPTEEEYLALCLIMLAR
 SGTGTRTGLTDATTSQQPADKKTAELPPVHKKEVATEQAEQSYKCSVCDKAFFSYQALGGHK
 ASHRKTTTATAASDDNNPSTSTSTGAVNISALNPTGRSHVCISICHKAFTGQALGGHKRRH
 YEGKLGGSNRDLGGGGGGHSGSVLTTSDGGASTHLDLNPASPELQLGLSIDCGRKS
 QLLPMVQEVESEPMAPKKPRLLFSLG

SEQ ID NO 18: Os_AF332876 Oryza sativa zinc finger transcription factor ZF1 mRNA, complete cds
 AATTCCGGCACGAGGCCACACAGCAACCAGCCAGCTGCCACACTAGCTTGAGGCGAGCGAGCG
 AAGCTTAGCTAGCGGATAGAACAAAGTCGTCGATCTGCTGCTTTGTGAATTGCGGTGG
 AAGCATGTCGAGCGCGTCGTCCATGGAAGCGCTCCACGCCGCGGTGCTCAAGGAGGAGCAGC
 AGCAGCACGAGGTGGAGGAGGCGACGGTGTGACGAGCAGCAGGCCACGAGCGGGGAGGAG
 GGCAGCACCTGCCCAAGGGTGGCGAAGCGGAAGCGGTGCGCCAGCGATCGGAGGA
 GGAGAACCTCGCGCTCGCCTCCTCATGCTGCCCGCGCCGCCACCCCGCTCCAGGC
 CGCCTCCGCTCTCGGCTCGCGCCCGGCCAGGTGCGGAGTTCAAGTGCTCCGCTGC
 GGCAAGTCCTCAGCTCCTACCAGGCCTCGGCGCCACAAGACGAGCCACCGGTCAAGCT
 GCCGACTCCGCCCGAGCTCCCGTCTGGCTCCGCCGCGCCCTGCTGCCCTCCCG
 CCGAGGACCGCGAGCCAGCCACGTATCCACCGCCGCGTCCCGACGGCATGACCAACAGA
 GTCCACAGGTGTCCATCTGCCAGAAGGAGTCCCCACCGGGCAGGCCTGGCGGACAA
 GAGGAAGCACTACGACGGTGGCTAGGCCTGGCGCCGGCATCTCAACCGAGCTCTGG
 CCACGGTGGCCGCCAGTCCGAGGGTGGGAAGCTCCGGCAACGCCAGTCCGCCACCCGGCG
 TTCGACCTCAACCTCCCAGCGTCCGGAGTTCTGTGGCGCCGTGCTCAAGGGCAAGAA
 GATGTGGGACGAGGAGGAGGAGGTCCAGAGCCCCCTGCCCTCAAGAACGCCGGCTTCTCA
 CCGCGTAATTCAGCAGCTGCACGGATCCGATCCGTCAAGAGTTTGTCTAGGGAGTGAATT
 CAGTCGAAACACACTATTGTTGATCGTTGTGCGCGTATTGTTAATTGTTCTGCTT
 TTGTACAGAGCAAGCGAGTGATACATAGCCATACATACAGTCATACAGATAAGGTCTAGCT
 CTTCTTGTTCTTGTAACTGGAACTGTACCTGTATCTTACACTTTGTTCTTGACA
 GTCATATATTGTAGACCAAAAAAAA

SEQ ID NO 19: Os_AF332876 Oryza sativa seedling zinc finger transcription factor ZF1, STZ ortholog, protein
 MSSASSMEALHAAVLKEEEQQQHEVEEATVVTSSSATSGEEGGHLPQGWAKRKRSPRQSEEE
 NLALCLLMLARGGHHRVQAPPPLSASAPPAGAEFKCSVCGKSFSSYQALGGHKTSRVKLP
 TPPAAPVLAPAPVAALLPSAEDREPATSSTAASSDGMTNRHRCSCIQKEFPTGQALGGHKR
 KHYDGGVGAGAGASSTELLATVAAESEVGSSGNGQSATRAFDLNLPAVPEFVWRPCSKGK
 WDEEEEVQSPLAFKKPRLLTA

FIGURE 3 (continued)

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SEQ ID NO 20: Ph_BAA05079_D26086.1 [Petunia x hybrida].
PETZFP4 zinc-finger protein gene

TTCACTCACCAAAACAACCTCTACCTCTTACTTGCACATTCAAATTCTTCATTACTA
 CTTATCTCTACTAATCTGATTGATTTAGTAAATCAAACAAGAGAATCTTTCAGTAATA
 CAAACAAAGAAAATTTCTCTATACTTGAATTGAGTTAGTAAGGCAAACAAGAAAATTC
ATGGCACTTGAAGCATTGAATTCTCCAACTACAACAACACCACCATTC
 AACGGGCTTAAGTACCTTGAGAGTGGACAAAAGGTAAAAGATCAAAAGGCAACGCAGCA
 TGGAACGACAGTGTACTGAAGAAGAGTATTGACACTTGTCTTATCATGCTAGCACGTAGC
 GATGGTTCTGTTAATAACTCACGGTCTCTACCCACCACCACTACCCACATCAGTTCCAGT
 AACGTCGCAAATAACGCGACGTTATTGGAACAGAAGAATTGTACAAGTGTCCGTTGTG
 GTAAAGGGTTGGGTCTTATCAAGCTTAGGTGGACATAAGCAAGTCACCGGAAACTTGTC
 AGCATGGGAGGAGATGAACAATCTACTACTTCCACTACTAAGCTAACGGGAACTAGTTC
 CGCTAACGTTAACGGTAACGGAAGAACTCACGAATGTTCAATTGTCACAAGTGCTTCTA
 CTGGACAAGCTTAGGTGGTCATAAAAGGTGCCACTATGACGGTGGTAACGTTAACGGTAAC
 GGAAGTGTAAAGTGTGGGTGACGTATCTGAAGGTGTGGGTCAACTATTAGTCATCACCG
 TGACTTTGACTTGAATATTCCCGCGTGTGCCGGAGTTTGCCGGATTGGTCCGGCGAGG
 ATGAGGGTGGAGAGTCCTCATCCAGCAAAGAAGTCAGGCTATCTCTTCCACCTAAACTGAA
 TTATTCAAAGGATTATAGAGGGAATTGATTGTTACAGGAAGATTATTAGGATTACAGA
 ATTTTTGTTGACTAGTTATGTAATAT

SEQ ID NO 21: Ph_BAA05079 [Petunia x hybrida] zinc-finger protein, STZ otholog, protein

MALEALNSPTTTPPSFQFENNGLKYLESWTKGRSKRQRSMERQCTEEYLALCLIMLARS
 DGSVNNRSRSLPPPLPPSVPVTSQINATLLEQKNLYKCSVCGKGFGSYQALGGHKASHRKLV
 SMGGDEQSTTSTTNVTGTSANVNGNGRTHECSICHKCFPTGQALGGHKRHYDGGNGNGN
 GSVSVGVTSSSEVGSTISHRDFDINIPALPEFWPGFSGEDEVESPHAKKSRLSLPPKLE
 LFKGL

SEQ ID NO 22: Ta_BAA03901 Triticum aestivum gene for zinc-finger protein WZF1, complete cds

ATGTCGTGTCGGCCATGGAAGCGCTCCACGCCCTGATCCCGGAGCAGCACAGCTGGACGT
 TGAGCGGGCTCGGGCTGTCAGCAGCGCCACCAGCGCGAGGAGAGCGGGCACGTGCTGCAGG
 GGTGGCCAAGAGGAAGCGATCGCGCCAGCGCTCCAGGAGAGAGAACCTCGCGCTCTGC
 CTCCTCATGCTCTCGCGCCGGCAAGCAGCGTGTCAAGTGTCCGCTCGCGCAAGTCCTTCAGCTC
 TGCGCCGGTGCCTGCCAGTTCAAGTGTCCGCTCGCGCAAGTCCTTCAGCTCCTACCAAG
 CGCTCGGAGGCCACAAGACGAGCCACCGGGTGAAGCAGCGTCTCCTCCCTGTATGCCGCT
 GCTGCCCACTCGTGGCCCTCCCGGCCGTCGCCGCATCTGCCGTCCGCCAGCCGGCAC
 GTCGTCACCGGCCGCTCCGACGGCGACCAACAGAGTCACAGGTGTCCATCTGCC
 AAAAGGAGTCCCGACTGGCGAGGGCTCGACCGAGCTCTGGCCGCCGGCCGAGTCTGAGGTGGG
 GTGGCGCCGCCGCTCGACCGAGCTCTGGCCGCCGGCCGAGTCTGAGGTGGG
 GAGCACCGGCAACGGGAGCTCCGCCGCCGGCCTCGACCTGAACATTCCGCCGTGCCGG
 AGTTCGTGTGGAGGCCGTGCCAACGGCAAGATGATGTGGGAGGACGATGAGGAGGTGCAG
 AGCCCCCTCGCCTCAAGAACGCTCGCTTCACCGCTTGA

FIGURE 3 (continued)

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SEQ ID NO 23: Ta_BAA03901_WZF1 *Triticum aestivum*, STZ ortholog, protein
 MSSSAMEALHALIPEQHQLDVEAAAATSSATSGEESGHVLQGWAKRKRSRRQRSEENLALC
 LLMLSRGGKQRVQAPQPESFAAPVPAEFKCSVCGKSFSYQALGGHKTSRVKQPSPPSDAA
 AAPlVALPAVAAILPSAEPATSSAASSDGATNRVHRC5ICQKEFPTGQALGGHKRHYDGG
 VGAAASSTELLAAAEESEVGSTGNSSAARAFDLNIPAVPEFWWRPCAKGKMMWEDDEEVQ
 SPLAFKKPRLLTA

SEQ ID NO 24: Ca AF539746 *Capsicum annum* zinc finger protein mRNA, complete cds
 AAAATCTCGCTACTTACTTACATCTTCTAGAATAGTCACTAGAACAGTAACCTTATACAA
 CGGATATCGATATGGCACTGAAGCTTGAATTCTCAACTGGTACACCAACTCCGCCACCG
 TTTCAATTGAGAGCAGCCACAGCTTCGATATATCGAAAATGGAGGAAGGGAAAGAG
 ATCTAAAAGGTACGCAGCATGGAGCACCAGCCTACTGAGGAAGAATACCTAGCGCTTGTGTT
 TGATCATGCTTGCACGTAGCGGTGGCTCCGTTAATCATCAACGATCTACCAACCGCCGGCT
 CCGGTGATGAAACTGCACCGCGCGTCGTATCATCGGCGGCGGAGGAGGAGAAGGAGAAGAT
 GGTGATAAGTGTGTTGTGGTAAGGGATTGGGCTTATCAAGCTTAGGTGGACACA
 AAGCTAGTCACCGGAAACTCGTACCCGGGAGATGATCAGTCACACTACCTCCACAACCACT
 AACGCAACCGGAAACAACACCTCCGTTAACGGCAACGGCAACAGAAGTGGAAAGGACTCACGA
 GTGTCGATTTGTACAAGTGTGTTCCACTGGACAAGCTTAGGTGGACACAAAAGGTGTC
 ACTACGACGGCGGTATCGTAACGGAAACGCTAACAGTGGCGTTAGTGCTAGCGTTGGAGTG
 ACGTCATCGGAGGGTGTGGGTCACAGTCAGTCACCGGGATTGCAACTTGAAACATTCCGGC
 GTTGCCGGAATTCTGGCTGGGATTGGTCCGGCAAGATGAGGTGGAGGAGTCCACATCCGG
 CGAAGAAATCGCGGTATGTTGCCAAATATGAATTATTCACATTAATGGGAAATT
 GATTGTTAGGATTTACTATTTGGTAGACAAAATTATACTATGTAAGTTTAATTTCATTG
 TGGGTGGGAGCAAATTTTAATTGGCTATAGACCTAGCTAGTTACTAATAGCAAAA
 TTCAATTGATTAAAAA

SEQ ID NO 25: Ca AF539746-*capsicum annum*, STZ ortholog, protein
 MALEALNSPTGPTPPPFQFESDGQQLRYIENWRKGKRSRSMEHQPTEEYLALCLIML
 ARSGGSVNHQRSLPPPAPVMKLHAPSSSSAAEEKEKMYKCSVCGKGFSYQALGGHKASH
 RKLVPGGDDQSTTSTTNATGTTSVNGNGNRSGRTHECSICHKCFPTGQALGGHKRHYDG
 GIGNGNANSGVSASVGTVSEGVSTVSHRDFDLNIPALPEFWLGFSGSEDEVESPHPAKKS
 RLCLPPKYELFQH

PARALOGS OF STZ IN ARABIDOPSIS THALIANA

SEQ ID NO 26: gi_18402298_ref_NM_112848.1 mRNA
 ACTTCACTCTCTAATTCTCTCTATCTCTCACCATATTCGCGATTTAAACTCTCAAC
 TTTCTCTCAAAATTCTGATCCTTGTAGCCAACAGTTAGAAGAAGATTGATCTGATCATGGC
 CCTCGAAGCGATGAAACACTCCAACCTTCTTCAACAGAATCGAAACGAAAGAAGATTGA
 TGAACGACGCCGTTTCATTGAGCCGTGGCTAAACGCAAACGCTCCAAACGTCAGCGTTCT
 CACAGCCCTTCTCGTCTTCCTCACCGCCTCGATCTGACCCAAATCCAGAATCAAGA
 TCTTACGGAAGAAGAGTATCTCGCTCTTGTCTCCTCATGCTCGTAAAGATCAACCGTCGC
 AAACGCGATTTCATCAACAGTCGCAATGTTAACGCCGCCAGAATCAAAGAACCTCCG
 TACAAGTGTAACTGTTGAAAAAGCGTTCTCTATCAGGCTTAGGCGGTACAAAGC
 AAGTCACCGAATCAAACCAACCGTAATCTCAACAAACCGCCGATGATTCAACAGCTCCGA

FIGURE 3 (continued)

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CCATCTCCATCGTCGCCGGAGAAAAACATCCGATTGCTGCCTCCGGAAAGATCCACGAGTGT
 TCAATCTGTATAAAGTGTTCGACGGGTCAAGCTTAAAGCGGTACACAAACGTTGTCACTA
 CGAAGGCAACCTCGCGCGGAGGAGGAGGAGGAGGAAAGCAAATCAATCAGTCACAGTGGAAAGCG
 TGTCGAGCACGGTATCGGAAGAAAGGAGCCACCGTGGATTCATCGATCTAAACCTACCGCG
 TTACCTGAACTCAGCCTCATCACAATCCAATCGTCACGAAGAGATCTGAGTCCGTTGAC
 CGGTAAGAACCGCTTTGTGACCGATCACGCCAACGACTCATCAAGAAAGAAGATTATCTT
 TAAAAATCTAACTACTCGACTATTAAATTCTGTGTGATTTTCGTTACAACCATAAGTTCA
 TTTTCATTTTTAGTTACAAATTAAATTGTTCTGATTGGATTGAATATTGGTATATTG
 TTAGGGGTTGATAC

SEQ ID NO 27: Translation of gi_18402298_ref_NM_112848.1
 MALEAMNTPTSSFTRIETKEDLMNDAVFIEPWLRKRKRSKRQSHSPSSSSPPRSRPKSQN
 QDLTEEEYLALCLLMLAKDQPSQTRFHQQSQSLTPPPESKNLPYKCNVCEKAFPSYQALGGH
 KASHRIKPPVTIISTTADDSTAPTISIVAGEKHPPIAASGKIHECSICHKVFPTGQALGGHKRC
 HYEGNLGGGGGGGSKSISHSGSVSSTVSEERSHRGFIDILNLPALPELSLHHNPIVDEEILSP
 LTGKKPLLLTDHDQVIKKEDLSLKI

SEQ ID NO 28: gi_30680473_ref_NM_120516.3_mRNA
 AAATCAAATCTTTCATTACAATTATCTTCTTCATAATTAGAAACTTAGTAGCTAGTCTT
 CAAGATAATGGCACTTGAAACTCTTACTTCTCCAAGATTATCTTCTCGATGCCGACTCTGT
 TTCAAGATTCACTAGGGTTTCATGGAAGCAAAGGCAAACGATCTAACGATCAAGATCT
 GAATTGACCGTCAGAGTCTCACGGAGGATGAATATATCGTTATGTCTCATGCTTCTTGC
 TCGCGACGGAGATAGAAACCGTGACCTTGACCTGCCTTCTTCTCGTCTTCACCTCCTCTGC
 TTCCTCCTTCTTCACTCCGATCTACAAGTGTAGCGTCTGTGACAAGCGTTTCGTCCTAC
 CAGGCTCTGGTGGACACAAGGCAAGTCACCGAAAAGCTTTCGCTTACTCAATCTGCCGG
 AGGAGATGAGCTGTCGACATCGTCGGCGATAACCACGTCGGTATATCCGGTGGCGGGGAG
 GAAGTGTGAAGTCGCACGTTGCTCTATCTGTCAAAATCGTTCGCCACCGGTCAAGCTCTC
 GGCAGGCCACAAACGGTGCCACTACGAAGGAAAAGACGGAGGCGGTGAGTAGTAGCGTGTC
 GAATTCTGAAGATGTGGGTCTACAAGGCCACGTCAGCAGTGGCCACCGTGGGTTGACCTCA
 ACATACCGCCGATACCGGAATTCTCGATGGTCAACGGAGACGAAGAGGTGATGAGTCCCTATG
 CCGCGAAGAAACTCCGGTTGACTTCCCGAGAAACCTAAACATAAAACCTAGGAAAACCT
 TTACAGAATTCAATTAGGAAATTGTTTACTGTATATAACAAATATCGATTGATTGAT
 GTTCTCTTCACTGAAAATTATGATTCTTGTATAATTGATGTTCTGAAAAAGATATTGAT
 AACTTTTATTGTTCACACGTATAAAATTGCTGGATACATCA

SEQ ID NO 29: Translation of gi_30680473_ref_NM_120516.3
 MALETLTSPRLSSPMPTLFQDSALGFHGSKGKRSKRSRSEFDRQSLTEDEYIALCLMLLARD
 GDRNRDLDLPSSSSSPPLLPLPTPIYKCSVCDKAFCSSYQALGGHKASHRKSFSLTQSAGGD
 ELSTSSAITTSGISGGGGGSVKSHVCSICHKSFATGQALGGHKRCHYEGKNGGGVSSSVNS
 EDVGSTSHVSSGHRGFDLNIPPIPEFSMVNGDEEVMSPPAKLRFDFPEKP

FIGURE 3 (continued)

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SEQ ID NO 30: *gi_30693252_ref_NM_114853.2_mRNA*

ATGGCTCTGACACTCTCAATTCTCCACCTCACCACCAACCACCGCTCCTCCTCCTT
 CCTCCGTTGCCTCGACGAAACCGAGCCGAAAACCTCGAATCATGGACCAAAAGAAAACGTA
 CAAAACGTACCGTATAGATCAACCAAACCTCCTCTCTGAAGAAGAGTATCTCGCTCTT
 TGCCTCCTTATGCTCGCTCGTGGCTCCTCGATCATCACTCTCACCCTCGGATCATCACTC
 TCTTCTCCACTGTCCGATCATCAGAAAAGATTACAAGTGTCCGTCGTGGCAAATCTTCC
 CGTCTTACCAAGCGTTAGTGGACACAAAACAAGTCACCGGAAACCGGTTAGTGTGATGTT
 AATAATAGTAACGGAACCGTTACTAATAACGGAATATTAGTAACGGTTAGTGGTCAAAG
 TGGGAAGACTCATAACTGCTCTATATGTTTAAGTCGTTCCCTCTGGTCAAGCATTGGGTG
 GTCACAAACGTTGTCACTATGATGGGTAACGTAACAGTAACTGGTACAATAGCCACAAG
 TTTGACCTAAATTACCGGCTGATCAAGTTAGTGATGAGACAATTGGAAAAGTCAACTCTC
 CGGTGAAGAACAAAGTCGGTGTGATTATTATTACCGATCGGGATTAGCTAG
 TGGTTGATCATTAGCTGAGTCGTAAATGAAAATGAT

SEQ ID NO 31: Translation of *gi_30693252_ref_NM_114853.2*

MALDTI NSPTTTTAPPPFLRCLDEPE NLESWTKRKRTKRHRIDQPNPPPSEEEYLAL
 CLLMLARGSSDHSSPPSDHHSLSPLSDHQKDYKCSVCGKSFPSYQALGGHKTSRKPVSVDV
 NNSNGTVTNNGNISNGLVGQSGKTHNCSICFKSFPSQALGGHKRCHYDGGNGNSNGDNSHK
 FDLNLPADQVSDETIGKSQLSGEETKSVL

SEQ ID NO 32: *gi_30694224_ref_NM_123683.2_mRNA*

AAATTTCTATAGCAATGGCGCTTGAGCTTAATTCAACCAAGATTGGTCGAGGATCCCTT
 AAGATTCAATGGCGTTGAGCAGTGGACCAAATGTAAGAAACGATCCAACGTTCGAGATCTG
 ATCTTCATCATAACCACCGTCTACTGAGGAAGAGTATCTAGCTTCTGTCTCATGCTCTT
 GCTCGGGATGGCGGCGATCTGACTCTGTGACGGTTGCGGAGAAGCCGAGTTATAAGTGTGG
 CGTTGTTACAAGACGTTTCTGCTTACCAAGCTCTCGCGGTATAAGCGAGCCACCGGA
 GCTTATACGGTGGTGGAGAGAATGATAATCGACACCATTCCACCGCGTGAATCTCACGTT
 TGTCGGTTGGAAATCTTCGCCACCGGTCAAGCTCTCGCGGCCACAAGCGGTGCCA
 CTACGATGGTGGCGTTCGAACTCGAAGGTGTGGGTCTACTAGCCACGTCAGCAGTAGTA
 GCCACCGTGGATTGACCTTAATATTACCGGTGCAGGGATTTCGCCGGACGACGAAGTG
 ATGAGTCCGATGGCGACTAAGAAGCTCGCCTGAAGTAAGTCTTTGTTGAAGACCTGGAAGT
 TTATCAAATGAAATATCAAATTCAAGGAACAGTTTGTGATTCTATTACCAAT
 ACACAATACGATTCAATTCC

SEQ ID NO 33: Translation of *gi_30694224_ref_NM_123683.2*

MALEALNSPRLVEDPLRFNGVEQWTKCKKRSKRSRSDLHHNNHRLTEEEYLAFCLMLLARDGG
 DLD SVTVAEKPSYKCGVCYKTFSSYQALGGHKASHRSLYGGGENDKSTPSTAVKSHVCVCG
 KSFATGQALGGHKRCHYDGGVNSEVGSTSHVSSSSHRGFDLNIIIPVQGFSPDDEVMS PMA
 TKKPRLK

FIGURE 3 (continued)

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SEQ ID NO 34: gi_30698307_ref_NM_126145.2_mRNA

CACACTCACTCTTCTTCATCTTCTTCTAAATAGCTGAAATCACATCTCACAGAAT
 TAAATCTTATGGCTCTCGAGACTCTCAATTCTCAACAGCTACCCACCACCGCTCGGCCTCTT
 CTCCGGTATCGTAAGAAATGGAGCCTGAGAATCTGAGCAATGGCTAAAGAAAAGAAC
 AAAACGTCAACGTTTGATCACGGTCATCAGAATCAAGAAACGAACAAGAACCTCCTCTG
 AAGAAGAGTATCTCGCTCTTGTCTCCTCATGCTCGCTCGTGGCTCCGCCGTACAATCTCCT
 CCTCTCCTCTACCGTCACGTGCGTCACCGTCCGATCACCGAGATTACAAGTGTACGGT
 CTGTGGAAAGTCCTTTCGTCATACCAAGCCTAGGTGGACACAAGACGAGTCACCGGAAAC
 CGACGAACACTAGTATCACTCCGGTAACCAAGAATGTCTAATAACAGTCACAGTAACAGC
 GGTCCGTTGTTATTAACGTTACCGTGAACACTGGTAACGGTGTAGTCAAAGCGGAAAGAT
 TCACACTTGCTCAATCTGTTCAAGTCGTTGCGCTGCTCAAGCCTTAGGTGGACACAAAC
 GGTGTCACTATGACGGTGGCAACACGGTAACGGTAACGGAAAGTAGCAGCAACAGCGTAGAA
 CTCGTCGCTGGTAGTGACGTAGCGATGTTGATAATGAGAGATGGTCCGAAGAAAGTGC
 CGGTGGCCACCGTGGATTGACCTAAACTTACCGTGTACAAGTCTCAGTGACGACTCTT
 AACGTTGACTGAGTTGAGGAAAAGTCAACTATCAAGCGAAGAAAGGGTTAGTGGACGGTG
 AAGATTAACGGTCGTTCTTCCAGTTGCTTCGGTTGAGCTGACTGGTCTGTAATGAAA
 ATGATGGAGTGGACTTGGCATTATTATTATTATTTTAAAAAGAAATGTTAATTGTTGTT
 GGATTTGTTATAGATAGAGGAAACAATTGGGATACACAAATATTTTTTTACAAAGA
 AAATAATAATGCAGAGATGGATGATGGATCGTACACGTTATTATATAGTGGACCATTCTG
 AATCGTGAATTATTATTATTGTTAGAAATTAAATTTCGT

SEQ ID NO 35: Translation of gi_30698307_ref_NM_126145.2

MALETLNSPTATT TARPLLRYREEMEPENLEQWAKRKRTKRQRFDHGHQNOETKNLPSEEE
 YLALCLLMLARGSAVQSPPLPLPSRASPSDHRDYKCTVCGKSFSYQALGGHKTSRKPTN
 TSITSGNQELSNSHNSNSGVVINVTVNTGNGVSQSGKIHTCSICFKSFASGQALGGHKRCH
 YDGGNNNGNGSSNSVEILVAGSDVSDVDNERWSEESAI GGHRGFDLNL PADQVSVTTS

OTHER GENES IN EVALUATION

SEQ ID NO 36: gi_12698881_ref_AF_332876.1_mRNA, 2xC2H2,
Oryza sativa

AATTCCGGCACGAGGCCACACAGCAACCAGCCAGCTGCCACACTAGCTTGAGGCGAGCGAGCG
 AAGCTTAGCTAGCGGATAGAACAAAGTCGATCTGCTGCTGCTTTGTGAATTGCGGTGG
 AAGCATGTCGAGCGCGTCGTCCATGGAAGCGCTCCACGCCGCGGTGCTCAAGGAGGAGCAGC
 AGCAGCACGAGGTGGAGGAGGCGACGGTCGTGACGAGCAGCAGGCCACGAGCGGGGAGGAG
 GGCAGACACCTGCCCAAGGGTGGCGAAGCGGAAGCGGTGCGCCGCCAGCGATCGGAGGA
 GGAGAACCTCGCGCTCGCTCCTCATGCTGCCCGCGCCACCACCGCGTCCAGCGC
 CGCCTCCGCTCGGCTCGGCCCGCCGGCAGGTGCGGAGTCAAGTCTCCGTCTGC
 GGCAAGTCCTCAGCTCTACCAGGGCGTCGGCGGCCACAAGACGAGCCACCGGGTCAAGCT
 GCCGACTCCGCCCGCAGCTCCGTTGGCTCCGCCCGTGCCTGCTGCCTTCCG
 CCGAGGACCGCGAGCCAGCCACGTATCCACCGCCGCGCTCCGACGGCATGACCAACAGA
 GTCCACAGGTGTTCCATCTGCCAGAAGGAGTCCCCACCGGGCAGGCCTGGCGGGCACAA
 GAGGAAGCACTACGACGGTGGCGTAGGCGCCGGCGCCGGCATCTCAACCGAGCTCCTGG
 CCACGGTGGCGCCGAGTCCGAGGTGGGAAGCTCCGGCAACGGCCAGTCCGCCACCCGGGCG
 TTCGACCTCAACCTCCGCCGTGCCGGAGTTCGTGTGGCGGCCGTGCTCCAAGGGCAAGAA
 GATGTGGGACGAGGAGGAGGAGGTCCAGAGCCCCCTGCCCTCAAGAAGCCCCGGCTTCTCA
 CCGCGTAATT CAGCAGCTGCACGGATCCGATCCGTCAAGAGTTTGTCTAGGGAGTCAAATT
 CAGTCGAAACACACTATT CGTTGATT CGTTGATTCGTTGATTCGCTATTGTTAATTGTTCTGCTT

FIGURE 3 (continued)

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TTGTACAGAGCAAGCGAGTGATACATAGCCATACATACAGTCATACAGATATAGGTCTAGCT
 CTTCCCTGGTCCTTGTAACACTGGAACGTACCTGTATCTTTACACTTGTCTTGACA
 GTCATATATTGTAGACCAAAAAAAAAAAAAAA

SEQ ID NO 37: *gi_12698882_ref_AAK01713.1, 2xC2H2, Oryza sativa*
 MSSASSMEALHAAVLKEEQQQHEVEEATVVTSSATSGEEGGHLPQGWAKRKRSSRRQRSEEE
 NLALCLLMLARGGHHRVQAPPILSASAPPAGAEFKCSVCGKSFSYQALGGHKTSHRVKLP
 TPPAAPVLAAPVAALLPSAEDREPATSSTAASSDGMTNRHRCSCICQKEFPTGQALGGHKR
 KHYDGGVGAGAGAGASSTELLATVAAESEVGSSGNGQSATRAFDLNLPAVPEFVWRPCSKGKKM
 WDEEEEVQSPLAFKKPRLITA

SEQ ID NO 38: *gi_6434215_ref_AL132966.1_region 116202 ... 116729, 2xC2H2, Arabidopsis thaliana*

ATGAAGAGAGACCGGTCGATTACGAAGAATCCATGAAGCATATAGACATAGTAGAAAGTCT
 AATGATGTTATCTCGAAGTTCTGTGTCAAACAAATCGATGTAAGCAATCTACCGGAAGCA
 AAACGAACCATAATAACCACTTCGAATGAAAACGTGTAACCGGAAATTGATTCCCTCCAA
 GCTCTGGAGGTATAGAGCTAGCCACAAGAACCTAACGCTGATCGTTGACCAAGAACAGGT
 GAAGCATCGTAACAAAGAGAATGATATGCATAAGTGTACAATTGCGATCAAATGTTGGGA
 CCGGTCAAGCTCTAGCGGTACATGAGAAAGCATAGGACGAGCATGATAACCGAGCAATCG
 ATTGTCCCTCTGTGGTTATTCCAGACCGGTTTTAACGTTGAGTAGCAGCAAGGAGAT
 CTTGGACTAAATCTAACTCCATTGGAAAATGATCTGTGTTAACCTTGGGAAGAATTGG
 TTCCACAAATTGATTGAAGTTGTGAATTAG

SEQ ID NO 39: *gi_6729511_ref_CAB67667.1, 2xC2H2, Arabidopsis thaliana*

MKRDRSDYEESMKHIDIVESLMMILSRSFVVKQIDVKQSTGSKTNHNNHFECKTCNRKFDSFQ
 ALGGHRASHKKPKLIVDQEQQVKHRNKENDMHKCTICDQMFGTQALGGHMRKRTSMITEQS
 IVPSVYVSRPVFNRCSSSKEILDNLTPLENDLVLIFGKNLVPQIDLKFN

SEQ ID NO 40: *ref_CA279020, 2xC2H2, sugar cane*
 CCTAACCGCATAGCTTTCAAATCAACAAGCCTGCCGTGACCGATCGATGGCCATCACC
 CACGACGACTACGTCTCCCTCTGCCCTATGGCGCTCGCAGCCGCGGGAGGCGGAGGCCAAGC
 TGGTTAACAAACCGAGTACGCTCTGAACACGGCTGCCCTGGACAGCGACGGCGCAAGAGTCG
 AGCTCCGCTTCCGGTCTCGTCTGTGGCAAGGCCTTCGCGTCGACCCAGGCACTGGCGGG
 CACAAGGCCAGCCACCGCAAGCCAGCCTGCTACAGGCACATGCGTCGTCCCTCAGCCGGAGG
 CGCGCGTCGTCGGTAACAATGACCTCGCCGTAGGCAGCAGTGGCAGGGGAGGCCACA
 GGTGCACGGTGTGCCATCGGAGCTTCGCGACGGNGCAAGCGCTGGCGGGCACAAGAGGTGC
 CATTACTGGGACGGGCTCTCGGTGTCGCTACCGCGTCGTCGGCGCCATGGGGTCCGGGTC
 GACCGTCAAGGGCTTGATCTGAATTGGTGCCGGTGCCTCCCGCATGGCCCAACGCTG
 CGACAAGGTGGGGAGAGGAGAANNAAGTCANAAAACCTGGCGGTCAAGAGAAGGCGGCTG
 CGGGTCCGTCTGGACCCCTAATTAAACGATTAGAAGTCTTTTAATGGATTGGGTGCTGGCGAA
 TTTTGAGAAGGTTGTAAGTTCTGAACCTGTTTAATGGATTGGGTGCTGGCGAA
 ATTTTAAACTGGATTAAATTGCGCTCACCTTTTTTATTACACCCTTTTT
 TTTTAGAAGAAGA

FIGURE 3 (continued)

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SEQ ID NO 41: *gi_18027011_ref_AF254447.1, 2xC2H2, Arabidopsis thaliana*

TCCTTCTCTCTCTCTCTCTCTCACCAGACTGATCCTTATTCCAATTCTTCACA
 GACTGGTTCAAGTCTAATCCTTTCAACCATTACCCCTAATTCCACTAACCCCTCTCCTCA
 TCCTCTCCTCCCTGTTACTCCTCCCTCTCCTCTTCTTCCCTCAATCCGGAGACCTCC
 GCCGTCCACCGCCGCCACCAACTCCCTCCCTCTCCTCCCTCTCCGAGAAGCCCTCCCTC
 CTCAGCCTCAGCCCCGCCAACAAACAAGACCCATCACAACCAGACCCCTTATTCA
 AGAACCCACCTTCACACCTCCATGGATGTCGACTACGATCATCACCATCAAGATGATCATATA
 ACCTCGATGACGATGACCATGACGTCACCGTTGCTCTCACATAGGCCTCCAAGCCCTAGT
 GCTCAAGAGATGCCCTTTGCTCATGATGTTCTTCTCCTCTCGAGGACCACTCA
 TCATCACGAGGACATGAATCACAAGAAAGACCTCGACCATGAGTACAGCCACGGAGCTGTCG
 GAGGAGGAGAAGATGACGATGAAGATTCACTGGCGGAGACGGCGCTGTAGAATCAGCAGA
 CTCAACAAGGTCATATTGGATCCCTACACCTCTCAGATTCTCATTGGCCCTACTCAGTT
 CTCATGTCCTGTTGCTCAAAACCTCAACAGATAAACATGCAGATGCATATGTGGG
 GACATGGATCACAATACAGAAAAGGACCTGAATCTAAGGGAAACACAACAGGAATG
 CTAAGGCTTCCCGTGTATTGCTGCGCCCCAGGCTGCGAACACATTGACCATCCAAGGGC
 AAAGCCTCTCAAAGACTTCAGAACCCCTCAAACACATTACAAGAGAAAACATGGGATCAAAC
 CTTTCATGTGTAGGAAATGGAAAGGCTTCGAGTCCGAGGGGACTGGAGAACACATGAG
 AAGAATTGTGGCAAACCTTGGTATTGCATATGTGGATCTGATTCAAGCACAAGAGATCTCT
 CAAAGATCACATCAAGGCTTTGGGAATGGTCACTGGAGCCTACGGAATTGATGGGTTTGATG
 AAGAAGATGAGCCTGCCTGAGGTAGAACAAATTAGACAAATGATCATGAGTCATGCAGTCT
 AAATAGCTTATATATATTACTATAAGTACTAAAGTAAATCGGTATATATATTAAAGAA
 ACCTAAATCTATGGACCAAGTTTGATGGAGGTAGGGCTTCAAAACTAAAGCTATATCAT
 CTAATTGATCATAGGAAAAAAATGAATCAAGAGCACTGGAAAATTAAATTGTATCTTA
 GCTTCCTAGTTAAATTATTGCAAGACAATGTAGCAGTCACCAATGAGGTTCCAACGGT
 TTATTCTATTGTATATTATTGTCATTAGCTCACCTTCGTTAATTGCAAGGACATAA
 CTTATAAAATGTTAAATTATG

SEQ ID NO 42: *At3g57670, 2xC2H2, Arabidopsis thaliana*

MTDPYSNFFTDWFKSNPFHYPNSSTNPSPHPLPPVPPSSFFFFPQSGDRLRRPPPPPTPPP
 SPPLREALPLLSLSPANKQDHHNHDHLIQEPPSTSMDVDYDHHDHHQDDHHNLDDDDHDVTV
 ALHIGLPSPSAQEMASLLMMSSSSSSRTTHHEDMNHKKLDHEYSHGAVGGGEDEDDDEDSV
 GGDGGCRISRLNKQYWIPTPSQILIGPTQFSCPVCFKTFNRYNMQMHWGHGSQYRKGP
 SLRGTOPTGMLRLPCYCCAPGCRNNIDHPRAKPLKFRTLQTHYKRKHGIKPFMCRKCGKAF
 AVRGDWRTHEKNCGKLWYCICGSDFKHRSILKDHIKAFGNHGAYGIDGFDEEDEPASEVEQ
 LDNDHESMQSK

SEQ ID NO 43: *gi_18676370_ref_AJ311810.2, 2xC2H2, Arabidopsis thaliana*

ATCTACACACTACTCACATCTCATCTCTCTAGCACATACCCATCAAACCATATAGAT
 ACGGTGTTTATTCTGATCTTCTCTTCTTCTCTCAGAGTCATGTCTAAT
 CCAGCTTGTGTTGAATCTCTCAACAACTGGATGTGACCATAATAGCTCAACTATTCCACTTC
 TCTCTCTACATTACAACACTCAGGTAGCTACTATTACTCTAAATACCAACCCCTAATT
 ACATTAATCATACTCATACCACTTCCACTTCCACTAACCTACCCCACTAAAGAGAAGCTCTT
 CCTCTCTTAGCTTAAGCCCCATAAGGCACCAAGAACACAAGACCAACACTATTCTATGGA
 CACCCATCAAATTAGCTCTCAAACCTTCTGATGATCCTCTGTGACTGTGGATCTCATC
 TAGGGTTACCAAACACTACGGTGTGGTGGAGAGCATTAGGAGCAATATTGCTCTGATGCAACC

FIGURE 3 (continued)

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ACGGACGAGCAAGATCAAGATCATGACCGAGGAGTAGAAGTCACAGTTGAGTCCCACCTTGA
 TGATGATGATGATCATCATGGAGATCTACACAGAGGTCACTACTATTGGATTCCACTCCTT
 CTCAGATTTGATTGGTCTACACAGTTCACTGTCTCTTGTCTCAAGACATTCAACAGA
 TACAACAACATGCAGATGCACATGTGGGGACACGGCTACAATACAGAAAGGGACAGAATC
 CTTAAGAGGAACCCAACCAACAGGAATGCTAAGACTACCATGTTCTGCTGTGCACCCGGTT
 GCAAGAACAAACATTGACCAACCCACAGAGCCAAGGCTCTTAAGGACTTCGAACCCCTCAAACA
 CATTACAAACGTAACATGGGTCTAAACCATTGCTGTCTATGTGGTAAGGCCTTGC
 AGTAAAAGGAGATTGGAGAACGCATGAGAAGAATTGTGGAAAGCTTGGTATTGCTCTTG
 GCTCGGATTTAACGACAAGAGGTGCGCTTAAGGACCATGTCAGGACCTTGGAAATGGTCAT
 GTTCCTGTGGGATTGATAGTTGGAGGAGATCATGAGGACTACTATGATGCTGCTCTGA
 TATCGAGCAATAAGATGATAGCAACAAACATGAGTGTAAATTAGGGTTTGTATTTC
 CTCTCATGCATTAGTTGATGTATGCACGTGTTAGTTGTCTTCGGATTTGT
 TATTTGTTTGAGCTGTTTTTTAATTACTAAGAAGTTAATTATCATCTAAAGATTT
 C

SEQ ID NO 44: gi_18376498_ref_CAC86167.1, 2xC2H2, *Arabidopsis thaliana*

MSNPACSNLFNNGCDHNSFNYSTSLSYIYNHSYSNTTNPNYINHTHTTSTSPNSPPLR
 EALPLLSLSPIRHQEQQDQHYFMDTHQISSSNFLDDPLVTVDLHLGLPNYGVGESIRSNIAP
 DATTDEQDQDHDRGVEVTVESHLDDDDHHGDLHHRGHYWIPTPSQILIGPTQFTCPLCFKT
 FNRYNNMQMHMWHGSQYRKGPESLRGTQPTGMLRLPCFCCAPGCKNNIDHPRAKPLKDFRT
 LQTHYKRKHGSKPFACRMCGKAFAVKGDWRTHEKNCGLWYCSCGSDFKHKRSLKDHVKAFG
 NGHVPKGIDSFGGDHEDYYDAASDIEQ

SEQ ID NO 45: gi_7798991_ref_AL355775.1_region 7957 ... 8451, 2xC2H2, *Arabidopsis thaliana*

ATGGTTGCGAGAAGTGAGGAAGTAGTGGAGATAGTGGAGATACGGCGCGAAATGTTGATGTT
 GTTATCAAGAGTTGGAGAATGCGGCGGAGGAGAGAAACGAGTTTCCGATGCAAGACTT
 GTCTAAAGAGTTTCGTCGTTCAAGCTTGGAGGTATCGTCAAGCCACAAGAAACTC
 ATTAACAGTAGCGATCCATCACTTCTGGATCCTGTCTAACAGAAAACCTAAACGGCGAC
 GTCTCATCCTGTCCGATATGTGGCGTGGAGTTCCGATGGGCAAGCTTGGTGGTACA
 TGAGGAGACATAGGAGTGAGAAGCTCACCAAGGCACGTTGGTACACGTTTTTACCG
 GAGACGACGACGGTGACGACTTGAAGAAACTCGAGTAGTGGGAAGAGAGTGGCTTGG
 CTTAGATTGCGATGGAGAGTTAGTCAATTGGAAGTTGGAGTTGGGAAGAACGATTCTGA

SEQ ID NO 46: gi_7798996_ref_CAB90935.1, 2xC2H2, *Arabidopsis thaliana*

MVARSEEVEIVEDTAKCLMLLSRVGECKGGGEKRVFRCKTCLKEFSSFQALGGHRASHKKL
 INSSDPSLLGSLSNKKTATSHPCPICGVFPMQALGGHMRHRSEKASPGTLVTRSFLP
 ETTTVTTLKSSSGKRVACLDLDSMESLVNWKLELGRITS

SEQ ID NO 47: gi_9755794_ref_AL391143.1_region 31730 ...

32938, 2xC2H2, *Arabidopsis thaliana*

ATGGGAAGACGAACATCAAGATCTCCATAAACCCATTAATGGAGCTTGTGAGACCTCAAGAT
 TACTCGGTACAGAAAGAACAGAAAAGTCTACGAACCAACAGCAAGATGTTACTTGTACT
 ATGGTCTAAGGGAAAACCTCGAAGAAGAAAACCCAGGAATCTCCGGAACCAATGAAGAAGATT
 TTGTTGATGCGAAGAATGTGGAAAGGGTTCGGTACGAGAAATTTAAGAATCATCG

FIGURE 3 (continued)

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CTCGATGATGCATTTATGCCGAACGAGAAGGTTGTGAAGAATCCTGATGACTCTGTCTC
 GTAGCCTTGGGTTGTGAAGAAGAAGAAAAGATCAAGACTTGGTAGGTCTGGGAAGACTTTA
 TTTACTACGTTCTGAACCGAGTTCTATTGGATGCGACTGATGAAGAATTAGAAGTGGC
 GGATTGTTGATTCTATTGCTAAGAGTGCTCCAAGGTTGAGACGAATTGAAAAGTCTT
 CTGAGGGAGTACGTGTTACTCCTGAAACACCTGAAAGTAGCTATGATTGGGTTGTTGCTC
 AACAAAGAAACCGAGAAAAGGTGGTGAATTGGAATCTGGGTTTAAGTAATGAGCAAAGACT
 TATGGAAGAAGGTTAGTAGTTATGGAACATCGAAAGAACCGAGCTAGCTTCTGAGAGACG
 AAAACAGATTGGATCAGCAGAAACGGAGAAAAGATGGTGAATTGAAATCCGGAATTGAGT
 AATGAGCAAAGACTGCTAGAAGAAGAGATTACTACTCCTGTGACATTCAAAGGTCCAGCGAG
 TTCCCTGAGACACAAGTGTGCTTGGATCGAAATGGAGGTGAATTGGCTCTGAGTTTGA
 GTAATGAGCAAACACTGATGGAAGAACATGGAAAGAACCGAGCTGAGTTCTTAGAAGATAAG
 CATGAATTGATCAGCGAAAATGCGAGAAGCTGGCGACTTGAATCTAGGTTTACAGAAT
 TGAGCTGGAGTAGGAGCTATGGAGTGTACTCTTCAGATACTGATATGCTCACGCAACTG
 ATAAGAAGAACGTTGAGCATCGATGCAGGTTGTCAACAAGATATTCTCGTTATCAAGCT
 CTAGGGGGTATCAGACGTTCATCGGATGAGCAAATGTAAGAACAGAACATGGCATAGA
 GGAATCAGTTGAACCCAGGATGACTCTGTGA

SEQ ID NO 48: gi_9755803_ref_CAC01747.1, 2xC2H2, Arabidopsis thaliana

MEDEHQDLHKPINGALRDLKITRSQKETEKSTNQQQDVTCYYGLRENSKKKTQESPEPMKKI
 LFRCEECGKGFRYEKYFKNHRSMMHLSNEKVEESLMTLSRSLGFVKKKRSRLGRSGKTL
 FTTFLEPSSIFDATDEELEVADCLILLSKSAPKVVDLKSLSAEAVRVPETPESSYDLGCLL
 NKKPRKGGELESVLSNEQRLMEEGFSYGTKEPASFLRDENRLDQQKRRKDFESESLLS
 NEQRLLEEEITTPVTFKGPASSLRHKCALDRNGGEFGPEFLSNEQTLMEETWKEPVSFLEDK
 HEFDQRMREAGDFESRFYRIELGVGAMECTSSTDMLTQSDKKNVEHRCRLCNKIFSSYQA
 LGGHQTFHRMSCKNNKNGIEESVEPRMTL

SEQ ID NO 49: gi_1418338_ref_X98678.1, 2xC2H2, Arabidopsis thaliana

CTTGTAGTTCACTCCACATAATAAACACCAAAGATTCTATTCTCTCCATAATTCGAA
 GTTTCTGAATTGGTTGTTCTGATTGTTCTGAATTGGGTTGGTCTCTTTCT
 TACTATATTGGATATGATGATGGTCAAGATGAGGTTGGGAGTGATCAGACGCAAATCATA
 AAAGGGAAACGTACGAAGCGACAAAGATCGTCTCGACGTTGTGGTACGGCGGCACAAAC
 AGTGACTTCAACAAGTTCATCGGCCGTGGAAGTGGAGGAGAAAGAGCTGTTCAGATGAAT
 ACAACTCGCGGGTTCTGTCCTCGGTGACTACTGATTGACGCAAGAAGAAGACATGGCG
 ATTTGTCATCATGTTAGCTCGTGGACAGTTCTCCATGCCGATCTCAAGAACTCGAG
 AAAAATTCACTCAGAAGATTCTCGCAGAATTCTAGTTCTATGTGTACGAGTGTAAACGT
 GTAACCGGACGTTCTCGTCTCCAAGCACTGGTGGACACAGAGCGAGCCACAAGAAGCCG
 AGGACGTCGACTGAGGAAAAGACTAGACTACCCCTGACGCCAACCAAGTCTAGTGCATCAGA
 AGAAGGGCAAACAGTCATTCAAAGTTCCGGCTCAGCCCTAGCTTACAGGCAAGTAACA
 TCATCAACAAGGCAAACAAAGTACACGAGTGTCCATCTCGCGTTCTGAGTTCACTCCGGG
 CAAGCTCTCGGTGGTCACATGAGGCGGCACAGGACAGCCGTAACCAAGCATTAGCCCCGTTGC
 AGCCACCGCAGAAGTAAGCAGAAACAGTACAGAGGAAGAGATTGAGATCAATATAGGCCGTT
 CGATGGAACAGCAGAGGAAATATCTACCGTTGGATCTTAAATCTACAGCACCAGGAGATGAT
 CTAAGAGAGTCCAAGTTCAAGGGATAGTATTCTCAGCAACACCAGCGTTAATAGATTGTCA
 TTACTAGTTGTTTTACTACATAATGATGAAATATTGTGAATTCTTACTTACT
 ACTATATTGTTGATCAAAAAAAAAAAAAAA

FIGURE 3 (continued)

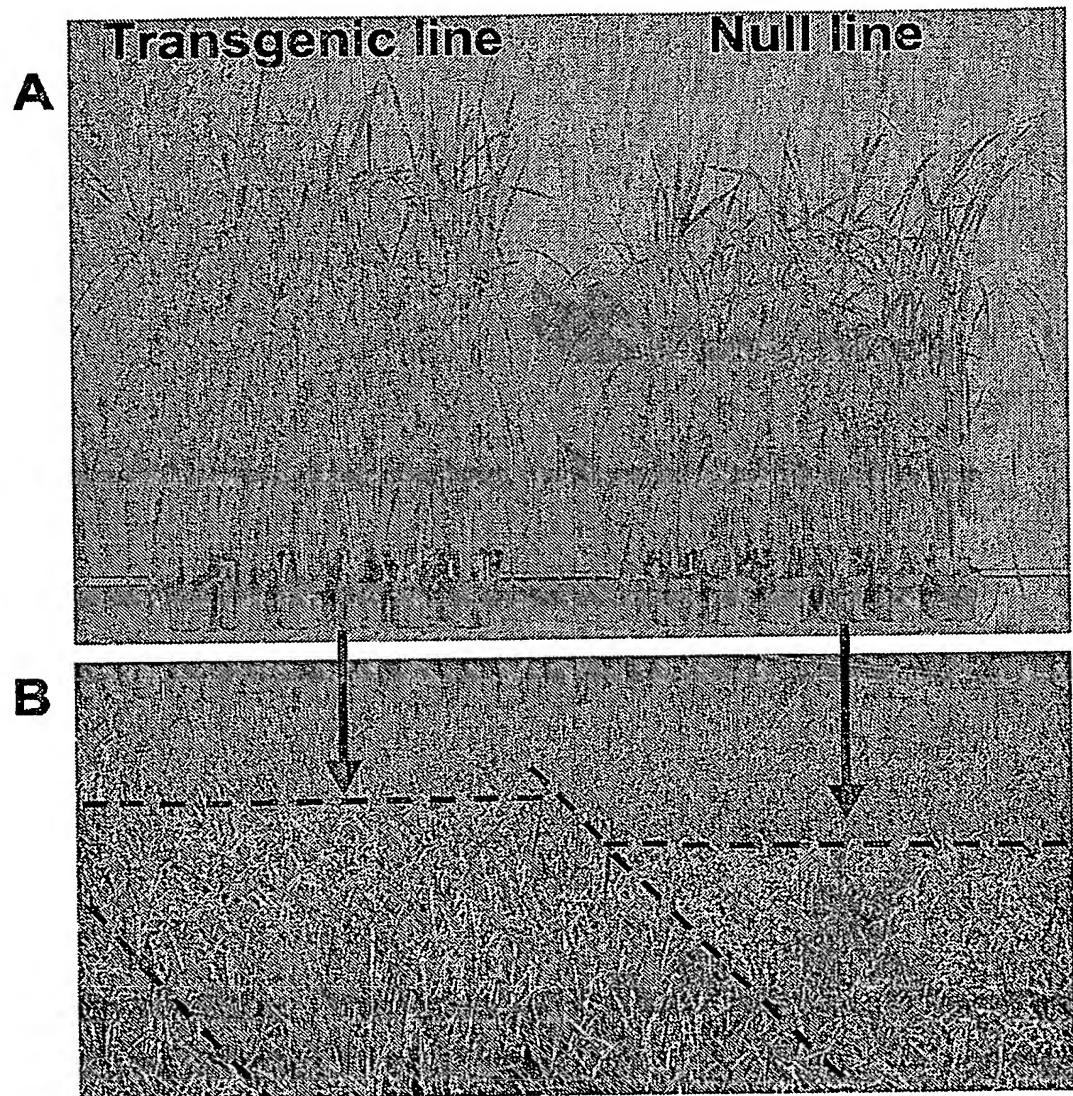
15/17

SEQ ID NO 50: gi_1418339_ref_CAA67236.1, 2xC2H2, Arabidopsis thaliana

MGQDEVGS DQTQI IKGKRTKRQRSSSTFVVTAA TTVTSTSSAGGS GGERAVSDEYNSAVSS
PVTTDCTQEEEDMAICLIMLARGTVLPSPDLKNSRKIHQKISSENSSFYVYECKTCNRTFSS
FQALGGHRASHKKPRTSTEKTRLPLTQPKSSASEEGQNSHFKVSGSALASQASNIINKANK
VHECSICGSEFTSGQALGGHMRRHRTAVTTISPVAATAEVSRNSTEEEIEINIGRSMEQQRK
YLPLDLNLPAPGDDLRESKFQGIVFSATPALIDCHY

FIGURE 3 (continued)

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**FIGURE 4**

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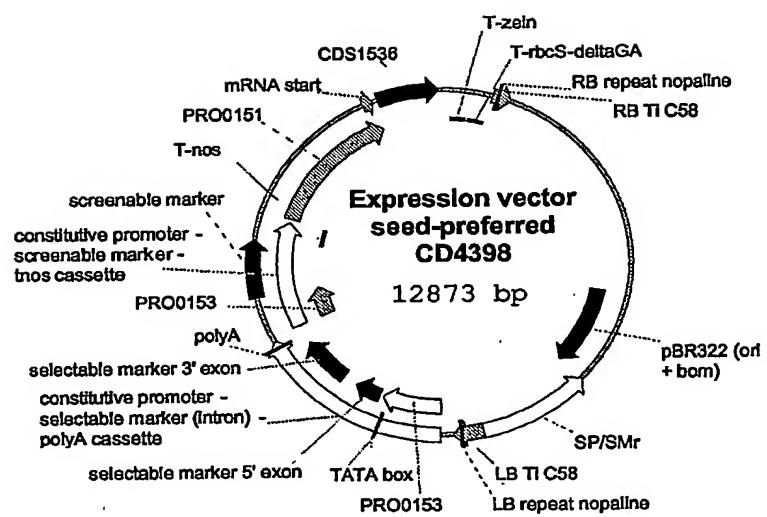


FIGURE 5